

SEQUENCE LISTING

<110> Robertson, David

Blakely, Randy D.

<120> GENETIC MUTATION UNDERLYING ORTHOSTATIC INTOLERANCE

5 AND DIAGNOSTIC AND THERAPEUTIC METHODS RELATING THERETO

<130> Attorney Docket No. 1242-27-2-2

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<150> 60/175,456

10 <151> 2000-01-11

<150> 60/173,682

<151> 1999-12-29

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<170> PatentIn Ver. 2.0

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<222> (1)..(1851)

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5 <301> Porzgen, P.

Bonisch, H.

Bruss, M.

<302> Molecular Cloning and Organization of the Coding Region

of the Human Norepinephrine Transporter Gene

10 <303> Biochem. Biophys. Res. Commun.

<304> 215

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<306> 1145-1150

<307> 1995

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Ala Asp Thr Gly Pro Glu Gln Pro Leu Arg Ala Arg Lys Thr Ala Glu  
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ctg ctg gtg gtg aag gag cgc aac ggc gtc cag tgc ctg ctg gcg ccc 144  
Leu Leu Val Val Lys Glu Arg Asn Gly Val Gln Cys Leu Leu Ala Pro  
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Leu Leu Val Val Lys Glu Arg Asn Gly Val Gln Cys Leu Leu Ala Pro  
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Arg Asp Gly Asp Ala Gln Pro Arg Glu Thr Trp Gly Lys Lys Ile Asp  
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Arg Phe Pro Tyr Leu Cys Tyr Lys Asn Gly Gly Ala Phe Leu Ile  
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Glu Leu Ala Leu Gly Gln Tyr Asn Arg Glu Gly Ala Ala Thr Val Trp  
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Lys Ile Cys Pro Phe Phe Lys Gly Val Gly Tyr Ala Val Ile Leu Ile  
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Tyr Tyr Leu Phe Ser Ser Phe Thr Leu Asn Leu Pro Trp Thr Asp Cys  
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Pro Ala Ala Glu Phe Tyr Glu Arg Gly Val Leu His Leu His Glu Ser  
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Val Leu Phe Val Leu Leu Val His Gly Val Thr Leu Pro Gly Ala Ser  
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	Ser Thr Phe Trp Ala Val Val Phe Phe Val Met Leu Leu Ala Leu Gly			
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	Ile Leu Phe Ala Val Leu Met Glu Ala Ile Gly Val Ser Trp Phe Tyr			
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	Pro Gly Leu Tyr Trp Arg Leu Cys Trp Lys Phe Val Ser Pro Ala Phe			
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	Ile Ala Leu Ser Ser Met Val Leu Val Pro Ile Tyr Val Ile Tyr Lys			
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Ala Asp Thr Gly Pro Glu Gln Pro Leu Arg Ala Arg Lys Thr Ala Glu  
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Leu Leu Val Val Lys Glu Arg Asn Gly Val Gln Cys Leu Leu Ala Pro  
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Arg Asp Gly Asp Ala Gln Pro Arg Glu Thr Trp Gly Lys Lys Ile Asp  
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Tyr Tyr Leu Phe Ser Ser Phe Thr Leu Asn Leu Pro Trp Thr Asp Cys  
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45 ggc cac acc tgg aac agc ccc aac tgt acc gac ccc aag ctc ctc aat 576  
Gly His Thr Trp Asn Ser Pro Asn Cys Thr Asp Pro Lys Leu Leu Asn  
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ggc tcc gtg ctt ggc aac cac acc aag tac tcc aag tac aag ttc acg 624  
Gly Ser Val Leu Gly Asn His Thr Lys Tyr Ser Lys Tyr Lys Phe Thr  
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	Pro Gly Leu Tyr Trp Arg Leu Cys Trp Lys Phe Val Ser Pro Ala Phe	
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	Leu Leu Phe Val Val Val Ser Ile Ile Asn Phe Lys Pro Leu Thr	
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	tac gac gac tac atc ttc ccg ccc tgg gcc aac tgg gtg ggg tgg ggc	1680
	Tyr Asp Asp Tyr Ile Phe Pro Pro Trp Ala Asn Trp Val Gly Trp Gly	
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	Ile Ala Leu Ser Ser Met Val Leu Val Pro Ile Tyr Val Ile Tyr Lys	
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	Phe Leu Ser Thr Gln Gly Ser Leu Trp Glu Arg Leu Ala Tyr Gly Ile	
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	acg cca gag aac gag cac cac ctg gtg gct cag agg gac atc aga cag	1824
	Thr Pro Glu Asn Glu His His Leu Val Ala Gln Arg Asp Ile Arg Gln	
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Ser Gly Ile His Asp Ile Gly Leu Pro Gln Trp Gln Leu Leu Leu Cys  
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Val Leu Phe Val Leu Leu Val His Gly Val Thr Leu Pro Gly Ala Ser  
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Thr Ser Phe Val Ser Gly Phe Ala Ile Phe Ser Ile Leu Gly Tyr Met  
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Ala Asp Thr Gly Pro Glu Gln Pro Leu Arg Ala Arg Lys Thr Ala Glu  
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Phe Leu Leu Ser Val Val Gly Phe Ala Val Asp Leu Ala Asn Val Trp  
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Glu Leu Ala Leu Gly Gln Tyr Asn Arg Glu Gly Ala Ala Thr Val Trp  
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	180	185	190	
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	195	200	205	
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	210	215	220	
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	225	230	235	240
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35	gct gga ttt gga gta ttg att gca ttt gcc agt tac aac aaa ttt gac Ala Gly Phe Gly Val Leu Ile Ala Phe Ala Ser Tyr Asn Lys Phe Asp	1008		
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40	aac aac tgt tac agg gat gcc ctg ctg acc agc agc atc aac tgt atc Asn Asn Cys Tyr Arg Asp Ala Leu Leu Thr Ser Ser Ile Asn Cys Ile	1056		
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	acc agc ttc gtc tct ggg ttc gcc atc ttc tcc atc ctt ggt tac atg Thr Ser Phe Val Ser Gly Phe Ala Ile Phe Ser Ile Leu Gly Tyr Met	1104		
	355	360	365	
45	gcc cat gaa cac aag gtc aac att gag gat gtg gcc aca gaa gga gct Ala His Glu His Lys Val Asn Ile Glu Asp Val Ala Thr Glu Gly Ala	1152		
	370	375	380	
	ggc cta gtg ttc atc ctg tat cca gag gcc att tct acc ctg tct gga Gly Leu Val Phe Ile Leu Tyr Pro Glu Ala Ile Ser Thr Leu Ser Gly	1200		
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Pro Tyr Thr Leu Phe Leu Ile Ile Ala Gly Met Pro Leu Phe Tyr Met  
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